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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/982,616

DATE: 02/19/2003

TIME: 14:37:11

Input Set : A:\09982616sequencelisting.txt
 Output Set: N:\CRF4\02192003\I982616.raw

4 <110> APPLICANT: Frances E. Lund
 5 Troy D. Randall
 6 Santiago Partida-Sanchez
 8 <120> TITLE OF INVENTION: CD38 MODULATED CHEMOTAXIS
 11 <130> FILE REFERENCE: AP33438 068443.0106
 13 <140> CURRENT APPLICATION NUMBER: 09/982,616
 14 <141> CURRENT FILING DATE: 2001-10-17
 16 <150> PRIOR APPLICATION NUMBER: 60/241,065
 17 <151> PRIOR FILING DATE: 2000-10-17
 19 <160> NUMBER OF SEQ ID NOS: 15
 21 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 23 <210> SEQ ID NO: 1
 24 <211> LENGTH: 1073
 25 <212> TYPE: DNA
 26 <213> ORGANISM: *Shistosoma mansoni*
 28 <400> SEQUENCE: 1
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 30 aattcaata atgatgaacg taatattgtt tcttacttta tcaaataattt ttgtctttaa 120
 31 ctctgcacaa catcaaataa acttacttag tgaaatagta caatcacat gtactcagtg 180
 32 gaagggtgaa catggagcta ctaatataag ttgttagtgat atctggatt cattgaaag 240
 33 cattttactt tcaactcata ctaatcagc atgtttatg aaatcagggt tattcgatga 300
 34 ttttggttat caattgtttg aatttggaca acaacaaca cagcgacacc acacaattca 360
 35 aacggaaacaa tactccatt ctcaagtgtt gaacatcatt cgttggatgt gtaaacgtct 420
 36 tgagttatgt cgttctctag aaactacatt tccaggatat ctgtttgtat aattgaattg 480
 37 gtgtatggc agtttaacag gcaacacaaa atacgggact gtatgtggat gcgattataa 540
 38 aagtaatgtt gttcatgcgt tctggaaag tgcttcggct gagttatgcc aggagacatc 600
 39 tggtaacatc tttgtggtag tgaatggctc ggtcaaagct ccatttaatg aaaataaaac 660
 40 ttttggaaaa atagaactac cattgttaaa acatcctcga gtacaacaat taacagtgaa 720
 41 attagttcat agtttggaaatgatgaaataa ccgacaaaaca tgtgaatcgtt ggagtctgca 780
 42 agaacttgca aacaagctga actctgtaca tattcctttt cgttgcattt acgatccttt 840
 43 agagttcaga cattatcaat gcattgaaaaa tcctggaaaaa caactatgtc agtttcagc 900
 44 ttcgacgagg tcaaacgtcg agacattact catactttt ccgcttagtca tttgtttaac 960
 45 tttttatact tccatgaatt gaaataactt ttcagaacta aactttgaac agagaaagag 1020
 46 aacaatgata ataaaggaat aggacattaa tgaaaaaaaaaaaaaaaaaaa aaa 1073
 48 <210> SEQ ID NO: 2
 49 <211> LENGTH: 353
 50 <212> TYPE: PRT
 51 <213> ORGANISM: *Shistosoma mansoni*
 53 <400> SEQUENCE: 2
 54 Glu Arg Thr Thr Tyr Ile Val Ile Ile Cys Ser Val Ile Phe His Asn
 55 1 5 10 15
 56 Leu Leu Ile Gln Ile Met Met Asn Val Ile Leu Phe Leu Thr Leu Ser
 57 20 25 30

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Input Set : A:\09982616sequencelisting.txt
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58 Asn Ile Phe Val Phe Asn Ser Ala Gln His Gln Ile Asn Leu Leu Ser
 59 35 40 45
 60 Glu Ile Val Gln Ser Arg Cys Thr Gln Trp Lys Val Glu His Gly Ala
 61 50 55 60
 62 Thr Asn Ile Ser Cys Ser Glu Ile Trp Asn Ser Phe Glu Ser Ile Leu
 63 65 70 75 80
 64 Leu Ser Thr His Thr Lys Ser Ala Cys Val Met Lys Ser Gly Leu Phe
 65 85 90 95
 66 Asp Asp Phe Val Tyr Gln Leu Phe Glu Leu Glu Gln Gln Gln Gln
 67 100 105 110
 68 Arg His His Thr Ile Gln Thr Glu Gln Tyr Phe His Ser Gln Val Met
 69 115 120 125
 70 Asn Ile Ile Arg Gly Met Cys Lys Arg Leu Gly Val Cys Arg Ser Leu
 71 130 135 140
 72 Glu Thr Thr Phe Pro Gly Tyr Leu Phe Asp Glu Leu Asn Trp Cys Asn
 73 145 150 155 160
 74 Gly Ser Leu Thr Gly Asn Thr Lys Tyr Gly Thr Val Cys Gly Cys Asp
 75 165 170 175
 76 Tyr Lys Ser Asn Val Val His Ala Phe Trp Gln Ser Ala Ser Ala Glu
 77 180 185 190
 78 Tyr Ala Arg Arg Ala Ser Gly Asn Ile Phe Val Val Leu Asn Gly Ser
 79 195 200 205
 80 -Val -Lys -Ala -Pro -Phe -Asn -Glu -Asn -Lys -Thr -Phe -Gly -Lys -Ile -Glu -Leu -
 81 210 215 220
 82 Pro Leu Leu Lys His Pro Arg Val Gln Gln Leu Thr Val Lys Leu Val
 83 225 230 235 240
 84 His Ser Leu Glu Asp Val Asn Asn Arg Gln Thr Cys Glu Ser Trp Ser
 85 245 250 255
 86 Leu Gln Glu Leu Ala Asn Lys Leu Asn Ser Val His Ile Pro Phe Arg
 87 260 265 270
 88 Cys Ile Asp Asp Pro Leu Glu Phe Arg His Tyr Gln Cys Ile Glu Asn
 89 275 280 285
 90 Pro Gly Lys Gln Leu Cys Gln Phe Ser Ala Ser Thr Arg Ser Asn Val
 91 290 295 300
 92 Glu Thr Leu Leu Ile Leu Phe Pro Leu Val Ile Cys Leu Thr Phe Tyr
 93 305 310 315 320
 94 Thr Ser Met Asn Asn Asn Phe Ser Glu Leu Asn Phe Glu Gln Arg Lys
 95 325 330 335
 96 Arg Thr Met Ile Ile Lys Glu Asp Ile Asn Glu Lys Lys Lys Lys Lys
 97 340 345 350
 98 Lys
 102 <210> SEQ ID NO: 3
 103 <211> LENGTH: 1050
 104 <212> TYPE: DNA
 105 <213> ORGANISM: Artificial Sequence
 107 <220> FEATURE:
 108 <223> OTHER INFORMATION: SM38 consensus sequence
 110 <400> SEQUENCE: 3
 111 ggaaagaacg tagacatata ttgttatata gatttgtca gttattttc acaatcttt 60

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Input Set : A:\09982616sequencelisting.txt
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112 aattcaaata atgatgaacg taatattgtt tcttacttta tcaaataattt ttgtctttaa 120
113 ctctgcacaa catcaaataa acttacttag taaaatagta caatcacat gtactcagt 180
114 gaagggtgaa catggagcta ctaatataag ttgttagttag atctggatt cattgaaag 240
115 cattttactt tcaactcata ctaaatcagc atgtgttatg aaatcagggt tattcgatga 300
116 tttgtttat caattgtttg aattggaaca acaacaacaa cagcgacacc acacaattca 360
117 aacggaacaa tacttccatt ctcaagtgtat gaacatcatt cgtggaatgt gtaaacgtct 420
118 tggagttatgt cggtctctag aaactacatt tccaggatat ctgtttgatg aattgaattt 480
119 gtgtatggc agttaaacag gcaacacaaa atacggact gtatgtgat gcgattataa 540
120 aagtaatgtt gttcatgcgt tctggcaaa tgcttcggct gagtatgcc 600
121 tggtaacatc tttgtggtag tgaatggctc ggtcaaagct ccatttaatg aaaataaaaac 660
122 tttggaaaaa atagaactac cattggttaa aacatcctcg agtacaacaa ttaacagtga 720
123 aattagttca tagttggaa gatgtaaata accgacaaac atgtgaatcg tggagtcgc 780
124 aagaacttgc aaacaagctg aactctgtac atattcctt tcgttgcatt gacgatcctt 840
125 tagagttcag acattatcaa tgcattgaaa atcctggcaa acaactatgt cagtttcag 900
126 cttcgacgag gtcaaacgtc gagacattac tcatacttt tccgctagtc atttggtaa 960
127 cttttatac ttccatgaat tgaatataact ttccagaact aaactttgaa cagagaaaga 1020
128 gaacaatgat aataaaggaa taggmcattta 1050
130 <210> SEQ ID NO: 4
131 <211> LENGTH: 473
132 <212> TYPE: DNA
133 <213> ORGANISM: *Shistosoma mansoni*
135 <400> SEQUENCE: 4
136 ggctgaggat -gccaggagag catctggtaa -catctttgtg- gtactgaatg gctcggtcaa 60
137 agctccattt aatgaaaata aaacttttgg aaaaatagaa ctaccattgt taaaacatcc 120
138 tcgagttacaa caattaacag tggaaatttagt tcatagtttg gaagatgtaa ataaccgaca 180
139 aacatgtgaa tcgtggagtc tgcaagaact tgcaaaacaag ctgaactctg tacatattcc 240
140 ttttcgttgc attgacgatc ctttagaggat cagacattat caatgcattt 300
141 caaacaacta tgcattttt cagttcgcac gaggtaaac gtcgagacat tactcatact 360
142 tttccgcta gtcatttttta taacttttta tactccatg aattgaaata actttcaga 420
143 actaaactttt gaacagagaa agagaacaat gataataaaag gaataggcca tta 473
145 <210> SEQ ID NO: 5
146 <211> LENGTH: 145
147 <212> TYPE: DNA
148 <213> ORGANISM: *Shistosoma mansoni*
150 <400> SEQUENCE: 5
151 atggaaaatc atggcaaaca actatgtcag ttttcagttt cgacgagggtc aaacgtcgag 60
152 acattactca tacttttcc gctagtcatt tttataacttc catgaattga 120
153 aataactttt cagaactaaa ctttg 145
155 <210> SEQ ID NO: 6
156 <211> LENGTH: 280
157 <212> TYPE: DNA
158 <213> ORGANISM: *Shistosoma mansoni*
160 <400> SEQUENCE: 6
161 ggagttatgtc gttctctaga aactacattt ccaggatatc tgtttgatga attgaattgg 60
162 tgcaatggca gtttaacagg caacacaaa tacggactg tatgtggatg cgattataaa 120
163 agtaatgtt ttcatgcgtt ctggcaaagt gcttcggctg agtatgcac gagagcatct 180
164 ggttaacatct ttgtggtagt gaatggctcg gtcaagatc catttaatga aaataaaaact 240
165 tttggaaaaa tagaactacc attggtaaa acatcctcg 280
167 <210> SEQ ID NO: 7

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Input Set : A:\09982616sequencelisting.txt
Output Set: N:\CRF4\02192003\I982616.raw

168 <211> LENGTH: 1049
 169 <212> TYPE: DNA
 170 <213> ORGANISM: *Shistosoma mansoni*
 172 <400> SEQUENCE: 7
 173 gaaaaagaacg tagacatata ttgttatata gattgttca gttatTTTC acaatCTTT 60
 174 aattcaaata atgatgaacg taatattgtt tcttacttta tcaaataattt ttgtcttaa 120
 175 ctctgcacaa catcaaataa acttacttag tgaataatgtt caatcacat gtactcagt 180
 176 gaagggttcaa catggagcta ctaatataag ttgttagttagt aaatcagggt tattcgatga 240
 177 cattttactt tcaactcata ctaaatcagc atgtgttatg aaatcagggt tattcgatga 300
 178 ttttgtttat caattgtttg aatttggaaaca acaacaacaa cagcgacacc acacaattca 360
 179 aacggaacaa tacttccatt ctcaagtgtt gaacatcatt cgtggatgt gttaacgtct 420
 180 tggagtagt cgttctctag aaactacatt tccaggatat ctgtttgatg aattgaattt 480
 181 gtgtatg c agtttaacag gcaacacaaa atacggact gtatgtggat gcgattataa 540
 182 aagtaatgtt gttcatgcgt tctggcaag tgcttcggct gagtatgcca ggagagcatc 600
 183 tggtaacatc tttgtggtag tgaatggctc ggtcaagct ccatttaatg aaaataaaaac 660
 184 ttttggaaaa atagaactac cattgttaaa acatcctcga gtacaacaat taacagtgaa 720
 185 attagttcat agtttggaaag atgttaataa ccgacaaaaca tgtgaatcgt ggagtctgca 780
 186 agaacttgc aacaagctga actctgtaca tattccttt cgttgcattt acgatcctt 840
 187 agagttcaga cattatcaat gcattggaaa tcctggcaaa caactatgtc agtttcagc 900
 188 ttgcacgagg tcaaacgtcg agacattact catactttt ccgctagtc tttgtttaac 960
 189 ttttataact tccatgaatt gaaataactt ttcagaacta aactttgaac agagaaagag 1020
 190 aacaatgata ataaaggaat aggacatta 1049
 192 <210>-SEQ-ID-NO:-8
 193 <211> LENGTH: 282
 194 <212> TYPE: PRT
 195 <213> ORGANISM: *Aplysia californica*
 197 <400> SEQUENCE: 8
 198 Met Ser Pro Val Ala Ile Val Ala Cys Val Cys Leu Ala Val Thr Leu
 199 1 5 10 15
 200 Thr Arg Ile Ser Pro Ser Glu Ala Ile Phe Pro Thr Pro Glu Leu Gln
 201 20 25 30
 202 Asn Val Phe Leu Gly Arg Cys Lys Asp Tyr Glu Ile Thr Arg Tyr Leu
 203 35 40 45
 204 Thr Ile Leu Pro Arg Val Lys Ser Asp Cys Arg Ala Leu Trp Thr Asn
 205 50 55 60
 206 Phe Phe Lys Ala Phe Ser Phe Lys Ala Pro Cys Asn Leu Asp Leu Gly
 207 65 70 75 80
 208 Ser Tyr Lys Asp Phe Phe Gln Arg Ala Gln Gln Thr Leu Pro Lys Asn
 209 85 90 95
 210 Lys Val Met Phe Trp Ser Gly Val Tyr Asp Glu Ala His Asp Phe Ala
 211 100 105 110
 212 Asp Asp Gly Arg Lys Tyr Ile Thr Leu Glu Asp Thr Leu Pro Gly Tyr
 213 115 120 125
 214 Met Leu Asn Ser Leu Val Trp Cys Gly Gln Arg Asp Lys Pro Gly Phe
 215 130 135 140
 216 Asn Gln Lys Val Cys Pro Asp Phe Lys Asp Cys Pro Val Gln Ala Arg
 217 145 150 155 160
 218 Glu Ser Phe Trp Gly Thr Ala Ser Ser Ser Tyr Ala His Ser Ala Glu
 219 165 170 175

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/982,616

DATE: 02/19/2003

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Input Set : A:\09982616sequencelisting.txt
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220 Gly Asp Val Thr Tyr Met Val Asp Gly Ser Asn Pro Lys Val Pro Ala
 221 180 185 190
 222 Tyr Arg Pro Asp Ser Phe Phe Gly Lys Tyr Glu Leu Pro Asn Leu Thr
 223 195 200 205
 224 Asn Lys Val Thr Lys Val Lys Val Ile Val Leu His Gln Leu Gly Gln
 225 210 215 220
 226 Lys Ile Ile Glu Arg Cys Gly Ala Gly Ser Leu Leu Asp Leu Glu Met
 227 225 230 235 240
 228 Val Val Lys Ala Lys Lys Phe Gly Phe Asp Cys Val Glu Asn Pro Lys
 229 245 250 255
 230 Ser Val Leu Phe Leu Leu Cys Ala Asp Asn Pro Asn Ala Arg Glu Cys
 231 260 265 270
 232 Gln Leu Ala Lys Arg Tyr Tyr Arg Ile Ala
 233 275 280
 236 <210> SEQ ID NO: 9
 237 <211> LENGTH: 300
 238 <212> TYPE: PRT
 239 <213> ORGANISM: Homo sapien
 241 <400> SEQUENCE: 9
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 243 1 5 10 15
 244 Arg Leu Ser Arg Arg Ala Gln Leu Cys Leu Gly Val Ser Ile Leu Val
 245 20 25 30
 246 Leu Ile Leu Val Val Val Leu Ala Val Val Val Pro Arg Trp Arg Gln
 247 35 40 45
 248 Gln Trp Ser Gly Pro Gly Thr Thr Lys Arg Phe Pro Glu Thr Val Leu
 249 50 55 60
 250 Ala Arg Cys Val Lys Tyr Thr Glu Ile His Pro Glu Met Arg His Val
 251 65 70 75 80
 252 Asp Cys Gln Ser Val Trp Asp Ala Phe Lys Gly Ala Phe Ile Ser Lys
 253 85 90 95
 254 His Pro Cys Asn Ile Thr Glu Glu Asp Tyr Gln Pro Leu Met Lys Leu
 255 100 105 110
 256 Gly Thr Gln Thr Val Pro Cys Asn Lys Ile Leu Leu Trp Ser Arg Ile
 257 115 120 125
 258 Lys Asp Leu Ala His Gln Phe Thr Gln Val Gln Arg Asp Met Phe Thr
 259 130 135 140
 260 Leu Glu Asp Thr Leu Leu Gly Tyr Leu Ala Asp Asp Leu Thr Trp Cys
 261 145 150 155 160
 262 Gly Glu Phe Asn Thr Ser Lys Ile Asn Tyr Gln Ser Cys Pro Asp Trp
 263 165 170 175
 264 Arg Lys Asp Cys Ser Asn Asn Pro Val Ser Val Phe Trp Lys Thr Val
 265 180 185 190
 266 Ser Arg Arg Phe Ala Glu Ala Ala Cys Asp Val Val His Val Met Leu
 267 195 200 205
 268 Asn Gly Ser Arg Ser Lys Ile Phe Asp Lys Asn Ser Thr Phe Gly Ser
 269 210 215 220
 270 Val Glu Val His Asn Leu Gln Pro Glu Lys Val Gln Thr Leu Glu Ala
 271 225 230 235 240

VERIFICATION SUMMARY

PATENT APPLICATION: **US/09/982,616**

DATE: 02/19/2003.

TIME: 14:37:12

Input Set : **A:\09982616sequencelisting.txt**
Output Set: **N:\CRF4\02192003\I982616.raw**